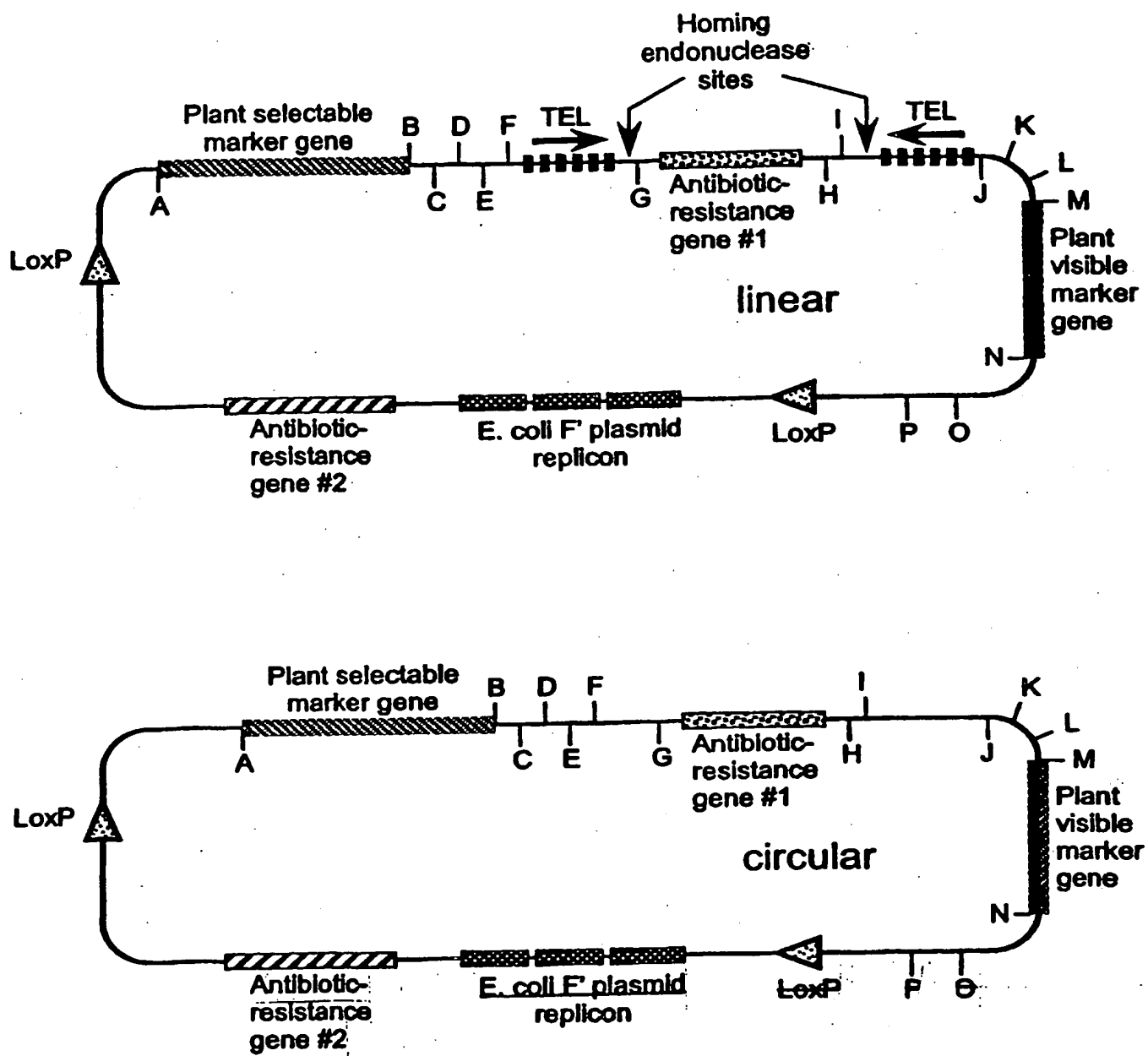


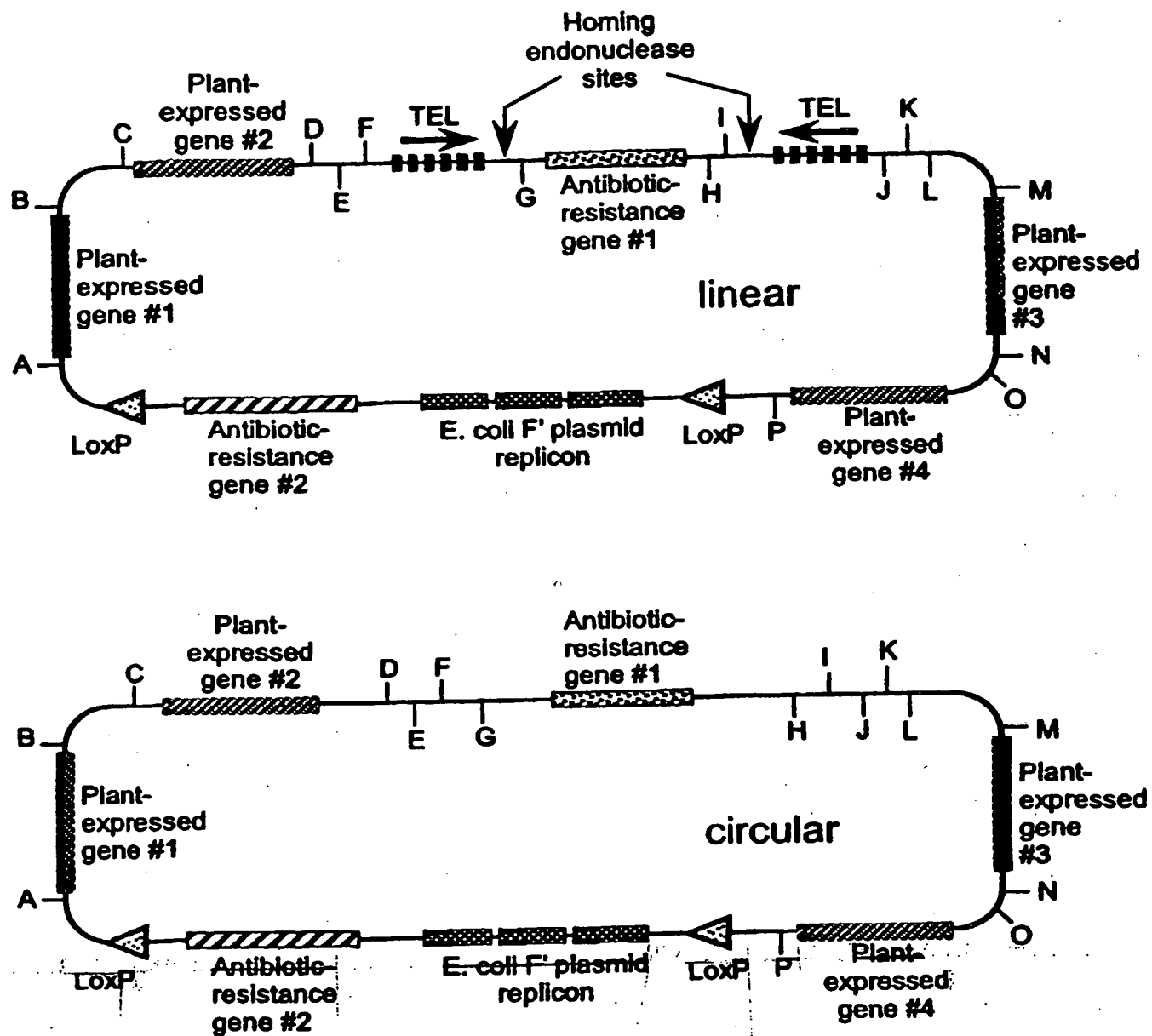
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Figure 1



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Figure 2



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Figure 3

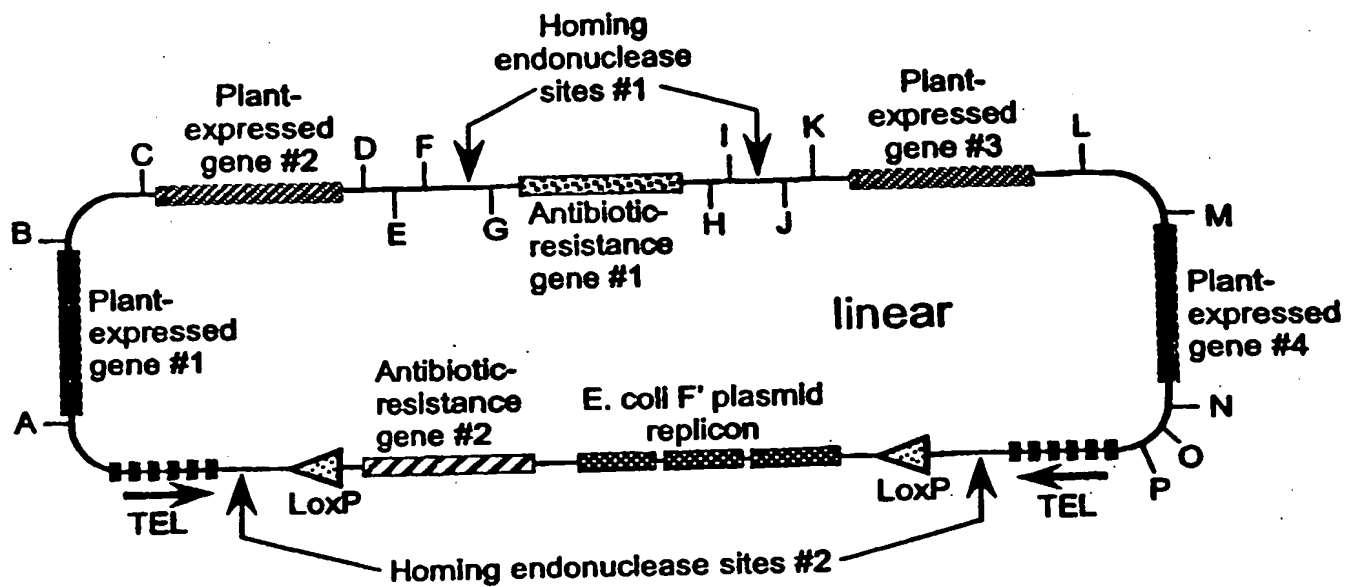


Figure 4

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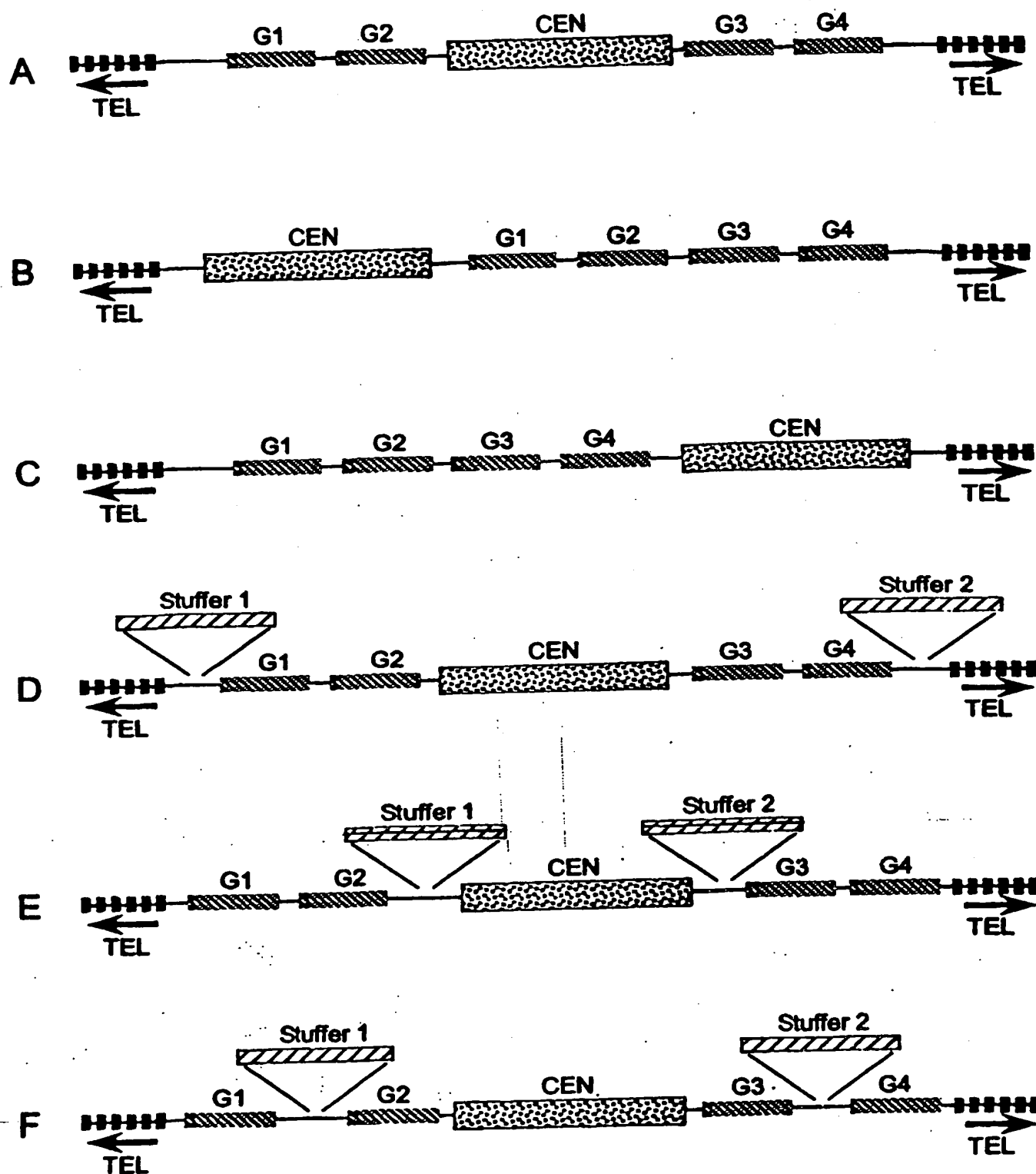


Figure 5

04	BB5R4-1	ASCTTSATTTGGATACATAAAAGTAGTGGAGAATCACCAGGAAGTTGAATAAA
05	BB5R4-1	AGCTTGATTTGGATACATAAAAGTGGTGGAGAATCACCAGGAAGTTGAATAAA
05	BB280R2-3	ACCTTCATTTGGATACATAAAAGTAGTGKAGAATCACCAGGAAGTTGAATAAA
04	BB5R4-1	TCTCATAGGAGTTAGGATGAAGAAGTTATCCCACTTTCAAATAAGGTGATCC
05	BB5R4-1	TCTCATAGGAGTTGGCATGAAGAAGTTATCCCMCTTTCAAATCAGGTGATTCT
05	BB280R2-3	TCTCATAGGAGTTAGGATGAAGAAGTTATCCCACTTTCAAATAAGGTGATCC
04	BB5R4-1	CAGTTTYCCTGTTTGGGAATATKANAACTTHTTTCGHCAATTCTADTCAAACCAG
05	BB5R4-1	CAGTTTCCCAGTTTGGGAATAGCACAGCTTCTTCGTCGTTCCAATCAAACCAG
05	BB280R2-3	CAGTTTYCCTGTTTGGGAATATGACAACTTCTTCGTCATTCTAATCAAACCAG
04	BB5R4-1	GATGAATCGCGATGTAARARVCY
05	BB5R4-1	GATGAATCTCTTTGTAAGA
05	BB280R2-3	GATGAATCKYGATGTWAGA

Figure 6

CrGm1	1..AAATTCAAAT	GGTCATAACT	TTTMACWCGG.....	30
CrGm2	1..AAATTCAAAC	GACAATAACT	TTTTACTCGG.....	30
SB12MC	1..AAAWTCAAAC	GACAATAACT	TTTKACTCGG.....	30
CrGm1	31..AKGTCCGATT	CAGGCGCATA	ATATATCGAG.....	60
CrGm2	31..ATGTCYGATT	GAGTCCCGTA	ATATATCGAG.....	60
SB12MC	31..ATGTCCGATT	GWGTCCCGTA	RTATATCGAG.....	60
CrGm1	61..ACGCTCGAAA	TTGAACAAYG	GAAGCTCTCG	AG..92
CrGm2	61..ACGCTCGAAA	TTGAATRYTG	-AAGCTCTGA	GC..92
SB12MC	61..ACGCTCGWAA	TTGAAACWG	-AAGCTCTRA	GM..92